

Honey Bee Mating Optimization with Nelder-Mead for Constrained Optimization, Integer Programming and Minimax Problems

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Received: 2 Sep. 2016, Revised: 27 Oct. 2016, Accepted: 2 Nov. 2016

Published online: 1 Jan. 2017

Abstract: In this article, we propose a new hybrid Honey Bee Mating Optimization (HBMO) algorithm with simplex Nelder-Mead method in order to solve constrained optimization, integer programming and minimax problems. We call the proposed algorithm a hybrid Honey Bee Mating Optimization (HBMONM) algorithm. In the the proposed algorithm, we combine HBMO algorithm with Nelder-Mead method in order to refine the best obtained solution from the standard HBMO algorithm. We perform several experiments on a wide variety of well known test functions, 6 constrained optimization problems, 7 integer programming and 7 minimax benchmark problems. We compare the performance of HBMONM against standard HBMO algorithm and Genetic Algorithm (GA). In the majority of tests, HBMONM is shown to converge faster, and reach a better solution than both HBMO and GA in reasonable time.

Keywords: Honey bee mating optimization, Nelder-Mead method, Genetic algorithm, Constrained optimization, Minimax problem, Integer programming, Swarm intelligence.

AMS subject classification: 90C10, 90C30, 90C47, 68T20.

1 Introduction

In recent years, several biological and natural processes have been influencing the methodologies in science and technology in an increasing manner. Among them, a number of swarm intelligence algorithms based on the behaviour of the bees have been presented [16]. These algorithms are divided, mainly, into two categories according to their behaviour in nature, their foraging behaviour and their mating behaviour. The most well known algorithm based on the marriage behaviour of bees is the Honey Bees Mating Optimization Algorithm (HBMO) that was presented in [1], [2] and simulates the mating process of the queen of the hive. Since then, it has been used on a number of different applications [3], [6], [9], [18], [19], [20], [30]

The (HBMO) algorithm belongs to a naturally inspired branch of algorithms called swarm intelligence (SI). SI are metaheuristic algorithms that consist of a decentralized population of individuals which interact locally with one another somewhat randomly. The local

interactions lead to a collective global intelligence that dictates the behavior of the population. Many SI algorithms are based on the behavior of animals or insects that tend to flock together, such as bat or ant colonies, herds of animals, and schools of fish such as the Particle Swarm Optimization (PSO) [16] and the cooperative behavior of bee colonies such as the Artificial Bee Colony (ABC) technique [14], the social foraging behavior of bacteria such as the Bacterial Foraging Optimization Algorithm (BFOA) [26], the simulation of the herding behavior of krill individuals such as the Krill Herd (KH) method [13], the mating behavior of firefly insects such as the Firefly (FF) method [32], [33] and the emulation of the lifestyle of cuckoo birds such as the Cuckoo Optimization Algorithm (COA) [27]. The HBMO algorithm aims to imitate the natural mating process of honey bees.

The goal of this work is to propose a new hybrid algorithm, namely, (HBMO) algorithm with simplex Nelder-Mead method in order to overcome the main

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drawbacks of the standard HBMO. In this paper, we propose a new hybrid algorithm, which is called simplex Honey Bee Mating Optimization (HBMONM) by combining HBMO and simplex Nelder Mead method in order to increase the exploration capability in the proposed algorithm and avoid stagnation and premature convergence in the population. Invoking the Nelder Mead method as a local search method in the final stage of the algorithm helps the proposed algorithm to accelerate the convergence and avoid performing iterations which do not improve the results. The HBMONM algorithm is tested on 6 constrained optimization problems, 7 integer programming and 7 minimax benchmark problems. The experimental results show that the proposed HBMONM is a promising algorithm and can obtain the optimal or near optimal solution for most of the tested function in reasonable time.

The organization of the paper is as follows. In Section ?? we present the basic algorithms such as the genetic algorithm (GA), the Nelder-Mead algorithm, and HBMO. In Section 3 we describe the proposed algorithm. In Section 4 we give the numerical experimental results for constrained optimization problems, integer programming, and minimax problems. Finally, in Section 5 we provide some concluding remarks and suggest future work.

2 The Basic Algorithms

2.1 Genetic Algorithm

The genetic algorithm (GA) is a metaheuristic algorithm that mimics natural selection and reproduction to find the global extrema. GA belongs to a larger class of evolutionary algorithms, which use biological mechanisms such as selection, reproduction, mutation, etc., to produce solutions to optimization problems. The main steps of GA are presented below:

Step 1. Randomly generate an initial population within the search space.

Step 2. Evaluate the fitness of each individual in the population.

Step 3. Choose parents according to their fitness.

Step 4. Use crossover operators on parents to produce offspring.

Step 5. Use mutation operators to alter the gene pool.

Step 6. Repeat steps 2–5 until termination criteria are met.

2.2 The Nelder-Mead algorithm

The Nelder-Mead (NM) algorithm is a derivative simplex method for finding minima for nonlinear functions [24]. The algorithm begins by creating a simplex of $n + 1$

vertices x_1, x_2, \dots, x_{n+1} ; where n is the dimension of the problem. The function is then evaluated at each vertex, and they are ordered according to their fitness such that x_1 and x_{n+1} correspond to the best and worst vertices respectively. At each iteration new vertices are computed to form a new simplex through four operations: reflection, expansion, contraction, and shrinkage. For each operation there is a corresponding scalar coefficient defined over a range: reflection $\rho > 0$, expansion $\chi > 0$, contraction $0 < \tau < 1$, and shrinkage $0 < \phi < 1$. The algorithm of Nelder-Mead is presented in Algorithm 2.

The main steps are presented below.

The Initial Simplex

Given an initial solution x , randomly generate n neighboring solutions to form the vertices of the simplex. the function of to be minimized is then evaluated at each point, and the vertices are reordered such that x_1 is the best point, and x_{n+1} is the worst point. The centroid of these points \bar{x} is calculated as:

$$\bar{x} = \frac{1}{n} \sum_{i=1}^n x_i \quad (1)$$

Reflection

The reflection process starts by computing the reflection point about the centroid $x_r = \bar{x} + \rho(\bar{x} - x_{n+1})$. If the reflected point $x_1 < x_r \leq x_n$, then the reflected point is accepted and replaces x_{n+1} . If $x_r < x_n$, then the algorithm proceeds to expansion.

$x_1 < x_r$, then the reflected point is accepted and replaces x_{n+1} . If x_r is less than x_n , then the algorithm proceeds to expansion.

Expansion

The expansion process starts by computing the expansion point $x_e = \bar{x} + \chi(x_r - \bar{x})$. If $x_e < x_1$, x_e replaces x_{n+1} ; otherwise x_r replaces x_{n+1} and the iteration terminates.

Contraction

If the reflected point $x_r > x_n$, the contraction process begins. There are two types of contractions: inside and outside. Which contraction is used depends on the comparison between x_r and x_{n+1} . If $x_n < x_r < x_{n+1}$, an outside contraction is performed according to $x_{oc} = \bar{x} + \tau(x_r - \bar{x})$. If $x_{oc} < x_r$ the contracted point is accepted and replaces the worst point. If $x_r > x_{n+1}$, then an inside contraction is performed according to $x_{ic} = \bar{x} + \tau(x_{n+1} - \bar{x})$. If $x_{ic} < x_{n+1}$ the contracted point is accepted and replaces the worst point. If either contraction is accepted, the iteration terminates.

Shrinkage

If no contraction point was accepted the shrink process executes by shrinking all points toward the best point to create a new simplex: x_1, x_2, \dots, x_{n+1} , where $x'_i = x_1 + \phi(x_i - x_1)$, $i = 2, \dots, n + 1$.

At the beginning of each iteration, the vertices are reordered, and the centroid recalculated.

Algorithm 1 The NM Algorithm

```

1: Let  $x_i$  denote the list of vertices in the current simplex,  $i = 1, \dots, n + 1$ .
2: 1. Order.
3: Order and relabel the vertices such that  $f(x_1)$  and  $f(x_{n+1})$  are the lowest and highest function values respectively.
4: while  $f(x_1) - f(x_{n+1}) > tolerance$  do
5:   2. Reflection.
6:   Compute the centroid of the simplex  $\bar{x} = \sum x_i/n, i = 1, \dots, n$ .
7:   Compute the reflection point  $x_r = \bar{x} + \rho(\bar{x} - x_{n+1})$ .
8:   if  $f(x_1) \leq f(x_r) < f(x_n)$  then
9:     Replace  $x_{n+1}$  with  $x_r$  and proceed to step 6.
10:  end if
11:  3. Expansion.
12:  if  $f(x_r) < f(x_1)$  then
13:    Compute the expansion point  $x_e = \bar{x} + \chi(x_r - \bar{x})$ .
14:    if  $f(x_e) < f(x_r)$  then
15:      Replace  $f(x_{n+1})$  with  $f(x_e)$  and proceed to step 6.
16:    else
17:      Replace  $f(x_{n+1})$  with  $f(x_r)$  and proceed to step 6.
18:    end if
19:  end if
20:  4. Contraction.
21:  if  $f_n \leq f(x_r) < f(x_{n+1})$  then
22:    Compute an outside contraction  $x_{oc} = \bar{x} + \tau(x_r - \bar{x})$ .
23:    if  $f_{oc} \leq f(x_r)$  then
24:      Replace  $x_{n+1}$  with  $x_{oc}$  and proceed to step 6.
25:    end if
26:  else
27:    Compute an inside contraction  $x_{ic} = \bar{x} + \tau(x_{n+1} - \bar{x})$ .
28:    if  $f(x_{ic}) \leq f(x_{n+1})$  then
29:      Replace  $x_{n+1}$  with  $x_{ic}$  and proceed to step 6.
30:    end if
31:  end if
32:  5. Shrinkage.
33:  Evaluate the  $n$  new vertices  $x'_i = x_1 + \phi(x_i - x_1), i = 2, \dots, n + 1$ .
34:  Replace vertices  $x_2, \dots, x_{n+1}$  with  $x'_2, \dots, x'_{n+1}$ .
35:  7. Reordering.
36:  Order and relabel the vertices such that  $f(x_1)$  and  $f(x_{n+1})$  are the lowest and highest function values, respectively.
37: end while

```

2.3 Honey Bee Mating Optimization Algorithm

The main steps in the original HBMO algorithm are presented below.

The Mating Flight

At the beginning of each mating flight, the speed and energy of the queen are randomly generated. A random drone is then generated and its fitness is evaluated. A successful mating between the queen and a drone is determined probabilistically through an annealing function as follows:

$$prob(Q, D) = e^{-\frac{\Delta(f)}{S(t)}}, \tag{2}$$

where $prob(Q, D)$ is the probability of the drones chromosome D being added to the spermatheca of the queen Q , $\Delta(f)$ is the difference between the fitness of the queen $f(Q)$ and the fitness of the drone $f(D)$, and $S(t)$ is the speed of the queen at time t .

A successful mating occurs if the value of $prob(Q, D)$ is greater than a randomly generated number in the range $[0, 1]$. If the mating is successful, then the drone's sperm is added to the queen's spermatheca.

After each attempted mating, the queen transitions to a new randomly generated drone, and the speed and energy of the queen decay according to the equations:

$$S(t + 1) = \alpha S(t), \tag{3}$$

$$E(t + 1) = E(t) - \beta, \tag{4}$$

$$\beta = 0.5 \frac{E(t_0)}{M}, \tag{5}$$

where α is the speed reduction variable, $E(t)$ is the energy of the queen at time t , β is the energy reduction after each transition, and M is the maximum number of mating flights.

The stopping criterion for each mating flight is reached when her spermatheca is full, or the speed or energy has reached its respective minima.

Breeding

After the mating flight is complete, random genes are selected from the queen's spermatheca and combined with the queen's genome using an intermediate crossover operator as follows:

$$x_i = q_i + a(d_i - q_i), \tag{6}$$

where x_i , q_i , and d_i are the chromosomes of the offspring, queen, and drone, and a is a scaling factor chosen uniformly at random over the interval $[-0.25, 1.25]$. $[0, 1]$ is a common range for a , but in our case, the larger variable range tended to produce better results. The intermediate crossover operator was chosen due to its promising performance with unconstrained problems[15].

HBMO is dissimilar to GA in creating offspring because in GA each offspring has two definite parents. Conversely, in HBMO every brood has the queen as the mother, but does not have a single drone as a father, and can have a genome consisting of a mixture of genes from the spermatheca.

Mutation of the Broods

Once all broods are created, workers are chosen according to their fitness using roulette wheel selection. A worker will apply a mutation to a brood and the brood's fitness is reevaluated. If the mutation worsens the brood's fitness, it is rejected and the brood's genome is left unchanged. Initially, the workers have equal chance to be chosen, however, after each iteration, the workers are

sorted according to the change in fitness of the broods, and the highest probability is assigned to the best worker. Gaussian, uniform, non-uniform, and boundary mutation operators are used to represent the workers.

Replacement of the Queen and choosing the Elites

The final step of the algorithm is to compare the most fit brood to the queen. If the brood is more fit than the queen, it becomes the queen for the next iteration. Otherwise, the queen remains unchanged for the next iteration. All remaining broods except for a user specified number of best broods are killed. The best broods are the elite of the population and are added to the queen's spermatheca for the next mating flight.

Algorithm 2 The HBMO Algorithm

```

1: Objective min or max  $f(x)$ ,  $x = (x_1, x_2, \dots, x_d)$ .
2: Randomly generate a population of  $n$  drone chromosomes with random solutions.
3: Find the best solution  $Q$  in the initial population.
4: while ( $t < MaxFlights$ ) do
5:   Randomly generate a speed  $s \in [0, MaxSpeed]$ .
6:   Randomly generate an energy  $E \in [0, 1]$ .
7:   while ( $s > MinSpeed$  &  $E > MinEnergy$  & spermatheca is not full) do
8:     Generate a drone  $D$  with random genes.
9:     Calculate mating probability  $prob(Q, D)$  from equation 1.
10:    if  $rand < prob(Q, D)$  then
11:      Add drone spermatheca.
12:    end if
13:     $s = \alpha s$ 
14:     $E = \beta E$ 
15:  end while
16:  for  $n = 1 : numOffspring$  do
17:    for  $i = 1 : d$  do
18:      Draw  $a$  from a uniform distribution in  $[0, 1]$ .
19:      Choose a random drone  $D$  in the spermatheca.
20:      Do intermediate crossover via  $x_{ni} = Q_i + a(D_i - Q_i)$ .
21:    end for
22:  end for
23:  Evaluate the fitness of broods  $X = x_{n \times i}$ .
24:  for  $m = 1 : numMutations$  do
25:    Select a random gene of a random brood.
26:    Select a worker (mutation operator) using roulette wheel selection.
27:    Apply mutation to the gene.
28:    Update the fitness of the brood.
29:    Update the fitness of the worker.
30:  end for
31:  Find the current best solution  $Q$ .
32:  Select elite solutions (excluding  $Q$ ) and add to next flight's spermatheca.
33: end while

```

3 The Proposed HBMONM Algorithm

In this section, we present the proposed HBMONM algorithm. The parameter settings used for the tests are shown in Table 2. The main steps of the algorithm are as follows.

Step 1. An initial population is generated randomly and each solution in the population has their fitness evaluated. The best solution is chosen as the queen.

Step 2. At the beginning of the queen's mating flight, her energy and speed are generated randomly. At each step of the mating flight a drone is generated randomly and mates with the queen according to an annealing function. The mating flight ends when the queen's energy or speed have reached their minimum value, or when her spermatheca is full.

Step 3. The queen's genes are randomly combined with genes from her spermatheca using an intermediate crossover operator to create broods.

Step 4. Workers are chosen according to their fitness to mutate the genes of the broods. Any improvements are kept.

Step 5. The broods are sorted according to their fitness, and the most fit brood is compared to the queen. If the brood is more fit, it becomes the queen for the next iteration. All remaining broods except for a user specified number of best broods are killed. The best broods are 'elites' that are automatically added to the queen's spermatheca for the next mating flight.

Step 6. Steps 2 through 5 are repeated until a set number of iterations have been completed. In order to increase the efficiency of the search, the NM algorithm as outlined in Algorithm 2 will be performed until the termination criterion are met.

The pseudocode for the HBMONM algorithm is outlined in Algorithm 3.

4 Results and Discussion

Twenty test functions from various categories were used to evaluate and compare the performance of the HBMO, HBMONM, and GA algorithms. The parameters for HBMONM, and GA are listed in Tables 1, 2 respectively. The values in [15] were used as a starting point for many of the parameter settings for both algorithms, but were modified to provide the best results. For certain parameters a range of values is given, as the value that provides the best computational result may be problem dependent.

4.1 Constrained optimization problems

Constrained optimization problems appear in many science, finance, operations research and engineering

Algorithm 3 The HBMONM Algorithm

```

1: Objective min or max  $f(x)$ ,  $x = (x_1, x_2, \dots, x_d)$ .
2: Randomly generate a population of  $n$  drone chromosomes with random solutions.
3: Find the best solution  $Q$  in the initial population.
4: while  $t < MaxFlights$  do
5:   Randomly generate a speed  $s \in [0, MaxSpeed]$ .
6:   Randomly generate an energy  $E \in [0, 1]$ .
7:   while ( $s > MinSpeed$  &  $E > MinEnergy$  & spermatheca is not full) do
8:     Generate a drone  $D$  with random genes.
9:     Calculate mating probability  $prob(Q, D)$  from equation 1.
10:    if  $rand < prob(Q, D)$  then
11:      Add drone spermatheca.
12:    end if
13:     $s = \alpha s$ 
14:     $E = \beta E$ 
15:  end while
16:  for  $n = 1 : numOffspring$  do
17:    for  $i = 1 : d$  do
18:      Draw  $a$  from a uniform distribution in  $[0, 1]$ .
19:      Choose a random drone  $D$  in the spermatheca.
20:      Do intermediate crossover via  $x_{ni} = Q_i + a(D_i - Q_i)$ .
21:    end for
22:  end for
23:  Evaluate the fitness of broods  $X = x_{n \times i}$ .
24:  for  $m = 1 : numMutations$  do
25:    Select a random gene of a random brood.
26:    Select a worker (mutation operator) using roulette wheel selection.
27:    Apply mutation to the gene.
28:    Update the fitness of the brood.
29:    Update the fitness of the worker.
30:  end for
31:  Find the current best solution  $Q$ .
32:  Select elite solutions (excluding  $Q$ ) and add to next flight's spermatheca.
33: end while
34: Rank the solutions and keep the best solution  $x_1$ .
35: Generate the remaining  $x_2, \dots, x_{n+1}$  vertices to be used in NM.
36: Apply the Nelder-Mead method, as shown in Algorithm 2, until termination criterion is met.

```

disciplines, such as pressure vessel design problem, welded beam design problem, reliability optimization problems, potential energy functions for protein design and so on. The general form of a constrained optimization is defined as follows:

$$\begin{aligned}
 &\text{Minimize } f(x), x = (x_1, x_2, \dots, x_n)^T, & (7) \\
 &\text{Subject to} \\
 &\quad g_i(x) \leq 0, i = 1, \dots, m \\
 &\quad h_j(x) = 0, j = 1, \dots, l \\
 &\quad x_l \leq x_i \leq x_u
 \end{aligned}$$

Table 1: Parameter settings for the HBMONM algorithm

Number of workers	4
Number of queens	1
Size of spermatheca	35–45
Number of broods	10–50
Maximum number of mating flights	10^4
Initial speed	$[0, 0.5-1]$
Speed reduction ratio	0.9
Minimum speed	$\frac{Speed}{1000}$
Initial energy	$[0, 1]$
Minimum energy	10^{-4}
Mutation rate	0.10–0.75
Number of elites	1–5
Tolerance	$10^{-7} - 10^{-3}$
ρ	1
χ	2
τ	0.5
ϕ	0.5

Table 2: Parameter settings for GA.

Population size	20
Selection function	Stochastic uniform
Fitness scaling	Rank
Maximum Generation	10^5
Crossover operator	Single point

where $f(x)$ is the objective function, x is a vector of n variables, $g_i(x) \leq 0$ are inequality constraints, $h_j(x) = 0$ are equality constraints, and x_l and x_u are variables bounds.

Evolutionary algorithms (EAs) have a number of advantages to solve constrained optimization, for example easy implementation, little information requirement for the problem to be solved, reliable and robust performance, etc. Due to those advantages, EAs have been successfully and broadly applied to solve COPs [5], [22], [23]. Many researchers have proposed various of EA-based constraint-handling techniques for real-parameter optimization problems which can be grouped as [5]: (1) hybrid methods; (2) separation of objectives and constraints; (3) special representations and operators; repair algorithms; (4) repair algorithms; (5) penalty functions.

The benchmark problems that were used are:

Test Problem 1 [10]. This problem is defined by

$$F_1(x) = (x_1 - 2)^2 + (x_2 - 1)^2 \tag{8}$$

subject to

$$x_1 = 2x_2 - 1, \quad \frac{x_1^2}{4} + x_2^2 - 1 \leq 0,$$

with

$$x_i \in [-100, 100], i = 1, 2.$$

The best known solution is $f^* = 1.3934651$.

Test Problem 2 [7]. This Problem is defined by

$$F_2(x) = (x_1 - 10)^3 + (x_2 - 20)^3, \quad (9)$$

subject to

$$\begin{aligned} 100 - (x_1 - 5)^2 - (x_2 - 20)^3 &\leq 0, \\ (x_1 - 6)^2 + (x_2 - 5)^2 - 82, 81 &\leq 0, \\ 13 \leq x_1 \leq 100, 0 \leq x_2 &\leq 100. \end{aligned}$$

The best known solution is $f^* = -6961.81381$.

Test Problem 3 [11]. This problem is defined by

$$\begin{aligned} F_3(x) = &(x_1 - 10)^2 + 5(x_2 - 12)^2 + x_3^4 + 3(x_4 - 11)^2 \\ &+ 10x_5^6 + 7x_6^2 + x_7^4 - 4x_6x_7 - 10x_6 - 8x_7, \end{aligned} \quad (10)$$

subject to

$$\begin{aligned} -127 + 2x_1^2 + 3x_2^4 + x_3 + 4x_4^2 + 5x_5 &\leq 0, \\ -282 + 7x_1 + 3x_2 + 10x_3^2 + x_4 - x_5 &\leq 0, \\ -196 + 23x_1 + x_2^2 + 6x_6^2 - 8x_7 &\leq 0, \\ 4x_1^2 + x_2^2 - 3x_1x_2 + 2x_3^2 + 5x_6 - 11x_7 &\leq 0, \end{aligned}$$

$$-10 \leq x_i \leq 10, \quad i = 1, \dots, 7.$$

The best known solution is $f^* = 680.370057$.

Test Problem 4 [11]. This problem is defined by

$$\begin{aligned} F_4(x) = &5.3578547x_3^2 + 0.8356891x_1x_5 + \\ &37.293239x_1 - 40792.141, \end{aligned} \quad (11)$$

subject to

$$\begin{aligned} 0 \leq &85.334407 + 0.0056858T_1 + \\ &T_2x_1x_4 - 0.0022053x_3x_5 \leq 92, \\ 90 \leq &80.51249 + 0.0071317x_2x_5 + \\ &0.0029955x_1x_2 + 0.0021813x_3^2 \leq 110, \\ 20 \leq &9.300961 + 0.0047026x_3x_5 + \\ &0.0012547x_1x_3 + 0.0019085x_3x_4 \leq 25, \end{aligned} \quad (12)$$

$$78 \leq x_1 \leq 102, 33 \leq x_2 \leq 45, 27 \leq x_i \leq 45, \quad i = 3, 4, 5.$$

Where $T_1 = x_2x_5$ and $T_2 = 0.0006262$. The best known solution is $f^* = -30665.538$.

Test Problem 5 [11]. This problem is defined exactly as Test Problem 4, except with

$$T_1 = x_2x_3, \quad T_2 = 0.00026.$$

The best known solution is unknown.

Test Problem 6 [21]. This problem is defined by

$$\begin{aligned} F_6(x) = &10.5x_1 - 7.5x_2 - 3.5x_3 - 2.5x_4 - \\ &1.5x_5 - 10x_6 - 0.5 \sum_{i=1}^5 x_i^2, \end{aligned} \quad (13)$$

subject to

$$\begin{aligned} 6x_1 + 3x_2 + 3x_3 + 2x_4 + x_5 - 6.5 &\leq 0, \\ 10x_1 + 10x_3 + x_6 &\leq 20 \\ 0 \leq x_i \leq 1, \quad i = 1, \dots, 5, &0 \leq x_6 \leq 50 \end{aligned}$$

The best known solution is $f^* = -213.0$.

For these test problems, the non-stationary penalty function employed in (Parsopoulos and Vrahatis 2002b) was used. The penalty function is defined in (Yang et al. 1997) as,

$$f(x) = F(x) + h(t)H(x) \quad (14)$$

where $F(x)$ is the original objective function of the constrained problem; $h(t)$ is a dynamically modified penalty value, where t is the current iteration number; and $H(x)$ is a penalty factor defined as

$$H(x) = \sum_{i=1}^m \theta(q_i(x))q_i(x)^{\gamma(q_i(x))} \quad (15)$$

where $q_i(x)$ is a relative violated function of the problems constraints, defined as $q_i(x) = \max\{0, g_i(x)\}$, $i = 1, \dots, m$, and $g_i(x)$ are the problem's constraints in the form $g_i(x) \leq 0$; $\theta(q_i(x))q_i(x)$ is a multi-stage assignment function (Homaifar et al. 1994); and $\gamma(q_i(x))$ is the power of the penalty function.

The parameters for the penalty function are problem dependent, using the values that provided the best results for the algorithms. The parameters are defined as

$$\gamma(q_i(x)) = \begin{cases} 1, & \text{if } q_i(x) < 0.01, \\ 2, & \text{otherwise,} \end{cases}$$

$$\theta(q_i(x)) = \begin{cases} 10, & \text{if } q_i(x) < 0.001, \\ 20, & \text{if } 0.001 \leq q_i(x) < 0.01, \\ 100, & \text{if } 0.1 \leq q_i(x) < 0.1, \\ 500, & \text{otherwise,} \end{cases}$$

and

$$h(t) = \begin{cases} \sqrt{t}, & \text{for Test Problem 1,} \\ t\sqrt{t}, & \text{otherwise.} \end{cases}$$

The problems' constraints in the form $g_i(x) \leq 0$ were only assumed violated if $g_i(x) > 10^{-5}$. In all test problems, HBMONM and GA were executed until 10^5 function evaluations were reached. The best feasible solution was then reported. For each test problem, 30 independent experiments were performed. The first experimental test was to compare HBMO to HBMONM

for constrained problems. The results of this are reported in Table 3. Figure 1 shows several examples where HBMONM reaches a lower function value faster than HBMO. Secondly, HBMONM was compared to GA, and the results are reported in Table 4. We can conclude from tables 3 and 4 that the combination of the standard HBMO algorithm with the NM algorithm can give improved results compared to both HBMO and GA.

Table 3: The mean function value after 10^5 function evaluations for the standard HBMO and hybrid HBMONM algorithms for constrained problems. The algorithm which displayed the best performance is in bold font.

Problem	HBMO	HBMONM
f_1	1.656	1.413
f_2	-6939.203	-6961.831
f_3	683.800	680.630
f_4	-30658.527	-30665.551
f_5	-31026.435	-31026.435
f_6	-212.996	-213.000

Table 4: The mean and the best solution found in all 30 runs for the constrained optimization problems. In parentheses is the sum of the violated constraints. The algorithm which exhibited the best performance is bolded.

Problem	Method	Mean Solution (Sum V.C.)	St.D.	Best Solution
f_1	HBMONM	1.413 (0.0001589)	0.0511	1.3934
	GA	2.0748 (0.000093)	0.5550	1.4746
f_2	HBMONM	-6961.831 (0.0002350)	0.0072	-6961.837
	GA	-6864.167 (0.0)	19.835	-6907.161
f_3	HBMONM	680.630 (0.0002237)	3.7406×10^{-6}	680.630
	GA	695.937 (0.0)	5.889	686.454
f_4	HBMONM	-30665.550 (0.0)	1.2878×10^{-6}	-30665.550
	GA	-30658.530 (0.0)	4.527	-30663.714
f_5	HBMONM	-31026.435 (0.0)	0.0	-31026.435
	GA	-31026.356 (0.0)	0.086	-31026.428
f_6	HBMONM	-213 (0.0)	4.4919×10^{-7}	-213
	GA	-212.997 (0.0)	0.003	-213.0

4.2 Minimax problems

The general form of the minimax problem is [34]

$$\min_x F(x), \tag{16}$$

where

$$F(x) = \max_{i=1, \dots, m} f_i(x), \tag{17}$$

with $f_i(x) : S \subset \mathbb{R}^n \rightarrow \mathbb{R}, i = 1, \dots, m$. Nonlinear programming problems of the form:

$$\begin{aligned} &\min_x F(x), \\ &g_i(x) \geq 0, \quad i = 2, \dots, m, \end{aligned}$$

can be solved as minimax problems of the form:

$$\min_x \max_{1 \leq i \leq m} f_i(x), \tag{18}$$

where

$$\begin{aligned} f_1(x) &= F(x), \\ f_i(x) &= F(x) - \alpha_i g_i(x), \\ \alpha_i &> 0, \end{aligned} \tag{19}$$

for $2 \leq i \leq m$. For sufficiently large values of α , it can be shown that nonlinear problems can be treated as minimax problems [4]. The benchmark problems that were used are:

Test Problem 7 [34]. This problem is defined by

$$\begin{aligned} &\min_x F_7(x), \\ &F_7(x) = \max\{f_i(x)\}, \quad i = 1, 2, 3, \\ &f_1(x) = x_1^2 + x_2^4, \\ &f_2(x) = (2 - x_1)^2 + (2 - x_2)^2, \\ &f_3(x) = 2e^{-(x_1+x_2)}. \end{aligned} \tag{20}$$

Test Problem 8 [34]. This nonlinear programming problem can be treated as a minimax problem according to (18) and (19). This problem is defined by

$$\begin{aligned} &F_8(x) = x_1^2 + x_2^2 + 2x_3^2 + x_4^2 - 5x_1 - 5x_2 - 21x_3 + 7x_4, \tag{21} \\ &g_2(x) = -x_1^2 - x_2^2 - x_3^3 - x_4^2 - x_1 + x_2 - x_3 + x_4 + 8, \\ &g_3(x) = -x_1^2 - 2x_2^2 - x_3^2 - 2x_4^2 + x_1 + x_4 + 10, \\ &g_4(x) = -x_1^2 - x_2^2 - x_3^2 - 2x_1 + x_2 + x_4 + 5. \end{aligned}$$

Test Problem 9 [34]. This nonlinear programming problem can be treated as a minimax problem according to (18) and (19). This problem is defined by

$$\begin{aligned} &F_9(x) = (x_1 - 10)^2 + 5(x_2 - 12)^2 + 3(x_4 - 11)^2 + x_3^4 \\ &\quad + 10x_5^6 + 7x_6^2 + x_7^4 - 4x_6x_7 - 10x_6 - 8x_7, \tag{22} \\ &g_2(x) = -2x_1^2 - 3x_3^4 - x_3 - 4x_4^2 - 5x_5 + 127, \\ &g_3(x) = -7x_1 - 3x_2 - 10x_3^2 - x_4 + x_5 + 282, \\ &g_4(x) = -23x_1 - x_2^2 - 6x_6^2 + 8x_7 + 196, \\ &g_5(x) = -4x_1^2 - x_2^2 + 3x_1x_2 - 2x_3^2 - 5x_6 + 11x_7. \end{aligned}$$

Test Problem 10 [29]. This problem is defined by

$$\begin{aligned} &\min_x F_{10}(x), \\ &F_{10}(x) = \max\{f_i(x)\}, \quad i = 1, 2, \end{aligned} \tag{23}$$

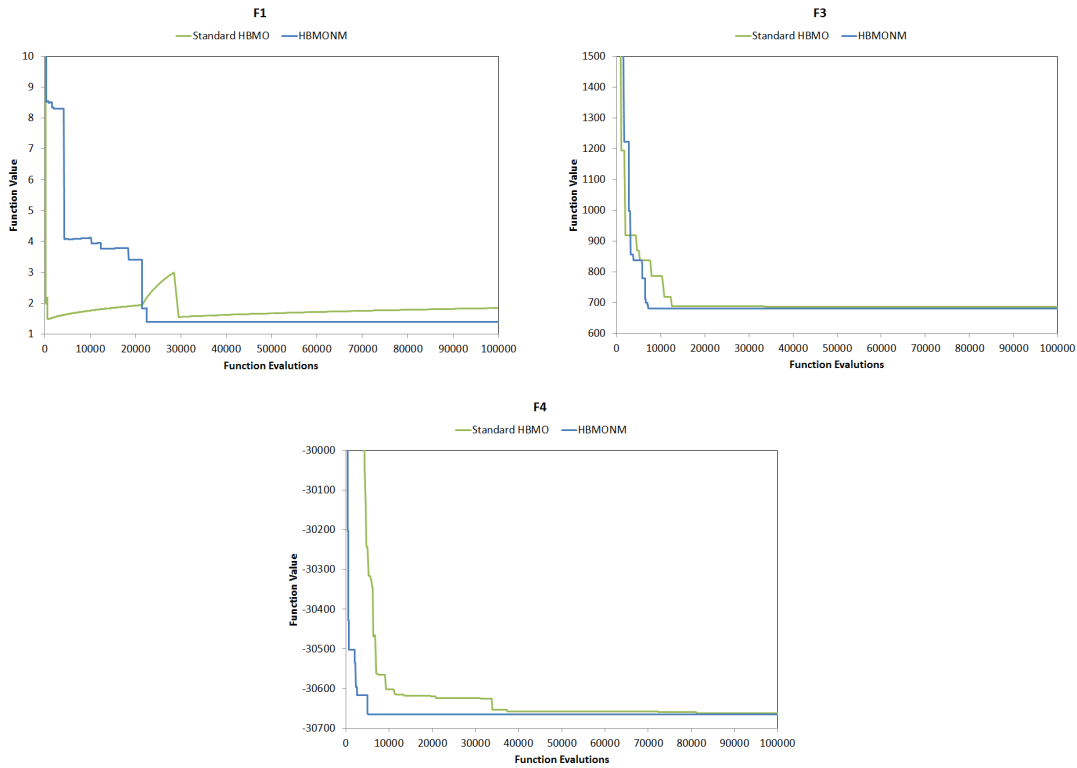


Fig. 1: The general performance of HBMONM and HBMO with constrained programming problems.

$$f_1(x) = |x_1 + 2x_2 - 7|,$$

$$f_2(x) = |2x_1 + x_2 - 5|.$$

$$f_i(x) = x_1 e^{(x_3 t_i)} + x_2 e^{(x_4 t_i)} - \frac{1}{1+t_i},$$

$$t_i = -0.5 + \frac{i-1}{20}.$$

Test Problem 11 [29]. This problem is defined by

$$\min_x F_{11}(x)$$

$$F_{11}(x) = \max\{f_i(x)\}, \tag{24}$$

$$f_i(x) = |x_i|, i = 1, \dots, 10.$$

Test Problem 12 [17]. This problem is defined by

$$\min_x F_{12}(x),$$

$$F_{12}(x) = \max\{f_i(x)\}, \tag{25}$$

$$f_1(x) = \left(x_1 - \sqrt{x_1^2 + x_2^2} \cos \sqrt{x_1^2 + x_2^2}\right)^2 + 0.005(x_1^2 + x_2^2),$$

$$f_2(x) = \left(x_2 - \sqrt{x_1^2 + x_2^2} \sin \sqrt{x_1^2 + x_2^2}\right)^2 + 0.005(x_1^2 + x_2^2).$$

Test Problem 13 [17]. This problem is defined by

$$\min_x F_{22}(x),$$

$$F_{22}(x) = \max\{|f_i(x)|\}, i = 1, \dots, 21, \tag{26}$$

For each test problem, 30 independent experiments were performed with $x \in [-50, 50]^n$, where n is the dimension of the problem. An experiment was considered successful only if it reached the desired error goal within 10^{-4} , and in less than 10^5 function evaluations. HBMO was first compared to HBMONM, and the results are presented in Table 5. The results from Table 5 show that HBMONM converges much faster than HBMO, and this is also clearly shown for a several test functions in figure 2. Next HBMONM was compared to GA, and the minimum, mean, maximum, and standard deviation of the required number of function evaluations are reported in Table 6. In all cases HBMONM performs better than both HBMO and GA.

4.3 Integer programming problems

An integer programming problem is a mathematical optimization problem in which all of the variables are restricted to be integers. The unconstrained integer programming problem can be defined as follows.

$$\min f(x), x \in S \subseteq \mathbb{Z}^n, \tag{27}$$

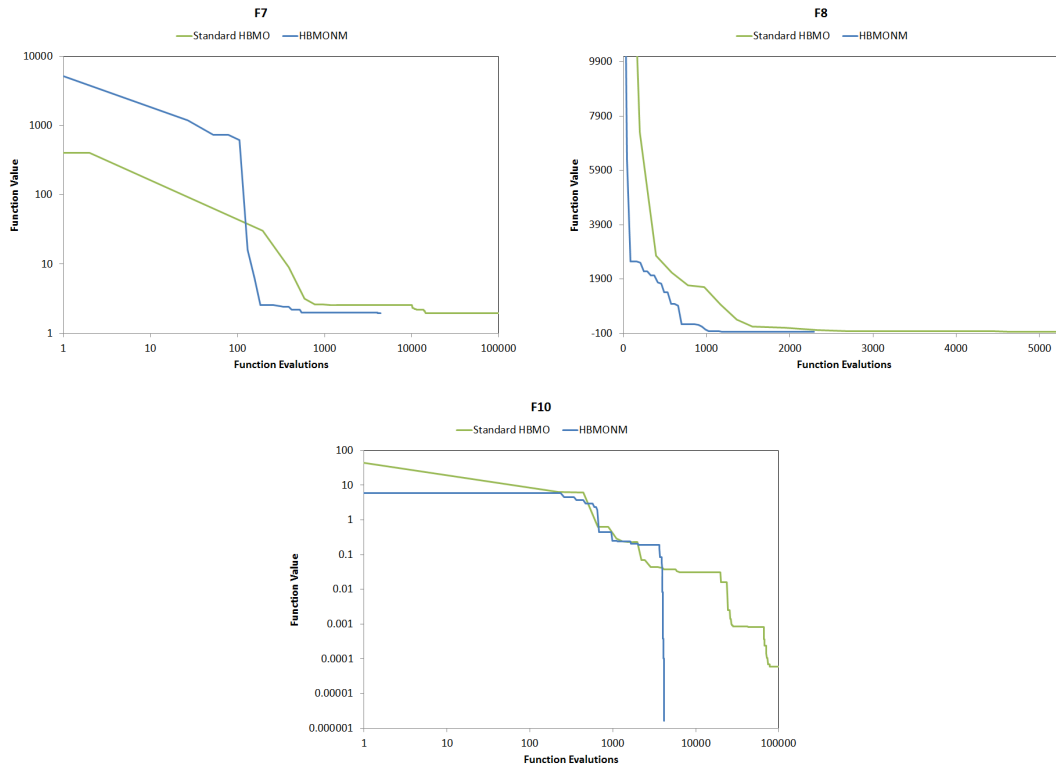


Fig. 2: The general performance of HBMONM and HBMO with minimax problems.

Table 5: The mean number of function evaluations to reach the desired error goal are shown for HBMO and HBMONM algorithms for minimax problems. If the desired error goal was not reached within 10^5 function evaluations, the experiment was stopped. The algorithm which displayed the best performance is in bold font.

Problem	HBMO	HBMONM
f_7	39073	658
f_8	78350	3714.73
f_9	10^5	6408.93
f_{10}	61358	1067.9
f_{11}	10^5	17225
f_{12}	58719	19060
f_{13}	65568	2013.7

Table 6: The results found in all 30 runs for the minimax optimization problems. The algorithm which exhibited the best performance is bolded.

Problem	Method	Mean	Min	Max	St.D.
f_7	HBMONM	658	99	1002	227.369
	GA	81960	75810	88040	2058.1
f_8	HBMONM	3714.73	1621	7478	1886.4
	GA	10^5	10^5	10^5	0
f_9	HBMONM	6408.93	3570	12044	240.551
	GA	10^5	10^5	10^5	0
f_{10}	HBMONM	1067.9	837	1503	153.167
	GA	87000	84375	92662	1025.741
f_{11}	HBMONM	17225	10578	22421	2635.1
	GA	10^5	10^5	10^5	0
f_{12}	HBMONM	19060	14561	25531	2550.6
	GA	38880	14260	71220	21134.7
f_{13}	HBMONM	2013.7	935	15345	2047.2
	GA	92212	90096	10^5	1434.4

where \mathbb{Z} is the set of integer variables, S is a not necessarily bounded set.

Now let us define the test problems.

Test Problem 14 [28]. This problem is defined by

$$F_{14}(x) = \|x\|_1 = |x_1| + \dots + |x_n|, \quad (28)$$

where the dimension of the problem is $n = 30$. The global minimum is $F_{14}(x^*) = 0$.

Test Problem 15 [28]. This problem is defined by

$$F_{15}(x) = x^T x = (x_1 \dots x_n) \begin{pmatrix} x_1 \\ \vdots \\ x_n \end{pmatrix}, \quad (29)$$

where the dimension of the problem is $n = 30$. The global minimum is $F_{15}(x^*) = 0$.

Test Problem 16 [8]. This problem is defined by

$$F_{16}(x) = -(15\ 27\ 36\ 18\ 12)x + x^T \begin{pmatrix} 35 & -20 & -10 & 32 & -10 \\ -20 & 40 & -6 & -31 & 32 \\ -10 & -6 & 11 & -6 & -10 \\ 32 & -31 & -6 & 38 & -20 \\ -10 & 32 & -10 & -20 & 31 \end{pmatrix} x, \quad (30)$$

The global minimum is $F_{16}(x^*) = -737$.

Test Problem 17 [8]. This problem is defined by

$$F_{17}(x) = (9x_1^2 + 2x_2^2 - 11)^2 + (3x_1 + 4x_2 - 7)^2, \quad (31)$$

The global minimum is $F_{17}(x^*) = 0$.

Test Problem 18 [8]. This problem is defined by

$$F_{18}(x) = (x_1 + 10x_2)^2 + 5(x_3 - x_4)^2 + (x_2 - 2x_3)^4 + 10(x_1 - x_4)^4, \quad (32)$$

The global minimum is $F_{18}(x^*) = 0$.

Test Problem 19 [8]. This problem is defined by

$$F_{19}(x) = 2x_1^2 + 3x_2^2 + 4x_1x_2 - 6x_1 - 3x_2, \quad (33)$$

The global minimum is $F_{19}(x^*) = -6$.

Test Problem 20 [8]. This problem is defined by

$$F_{20}(x) = -3803.84 - 138.08x_1 - 232.92x_2 + 123.08x_1^2 + 203.64x_2^2 + 182.25x_1x_2, \quad (34)$$

The global minimum is $F_{20}(x^*) = -3833.12$.

For each test problem, 30 independent experiments were performed with $x \in [-100, 100]^n$. An experiment was considered successful if the global minimum was found within 10^5 function evaluations. For both algorithms, the variables were only rounded to the nearest integer for function evaluations, and considered real numbers for all other purposes. The mean number of function evaluations required to reach the global minimum are compared for HBMO and HBMONM. The results are reported in Table 7. Next, HBMONM was compared to GA for the same set of problems, and the mean, minimum, maximum, and standard deviation of the required function evaluations are presented in Table 8.

The results of Table 7 show that HBMONM performs better on the majority of the test problems. Comparison of

the two algorithms is shown in Figure 3. Although HBMONM performed better than HBMO, Table 8 shows that GA performed better than both HBMO and HBMONM in the majority of integer programming problems.

Table 7: The mean number of function evaluations to reach the global minimum are shown for HBMO and HBMONM algorithms for integer problems. If the minimum was not reached within 10^5 function evaluations, the experiment was stopped. The algorithm which displayed the best performance is in bold font.

Problem	HBMO	HBMONM
f_{14}	31318.533	14376
f_{15}	15543.4	13354
f_{16}	63457	14776
f_{17}	187.4	652.24
f_{18}	10237.6	7454.1
f_{19}	484.67	525.7
f_{20}	661	1300.6

Table 8: The mean, min, and max number of function evaluations found in all 30 runs for the integer optimization problems. The algorithm which exhibited the best performance is bolded.

Problem	Method	Mean	Min	Max	St.D.	Successes
f_{14}	HBMONM	14376	9419	34377	6716.6	30/30
	GA	33364	17700	59040	12850	30/30
f_{15}	HBMONM	13354	6178	27861	4988.5	30/30
	GA	39637.143	12160	83520	22154.3	28/30
f_{16}	HBMONM	14776	2816	53191	10565	30/30
	GA	10474.667	4400	22380	5354.363	30/30
f_{17}	HBMONM	652.24	62	5798	1174	30/30
	GA	520	180	960	238.22	30/30
f_{18}	HBMONM	7454.1	4312	19410	2953.8	30/30
	GA	5006.67	2480	10480	1921.37	30/30
f_{19}	HBMONM	525.7	13	2139	582.78	30/30
	GA	841.33	140	2000	496.52	30/30
f_{20}	HBMONM	1306	78	6583	1331.6	30/30
	GA	620	240	2000	404.84	30/30

5 Conclusion

The performance of the HBMO algorithm for constrained, minimax, and integer optimization problems was compared to a new proposed hybrid HBMONM algorithm. Through their performance on numerous widely used, well known test functions from each category, it has been shown that HBMONM consistently performs better than the standard HBMO algorithm for the majority of problems. HBMONM was then compared with GA and it was shown that HBMONM performs better than GA for all constrained and minimax problems tested, however, GA performs better on the majority of

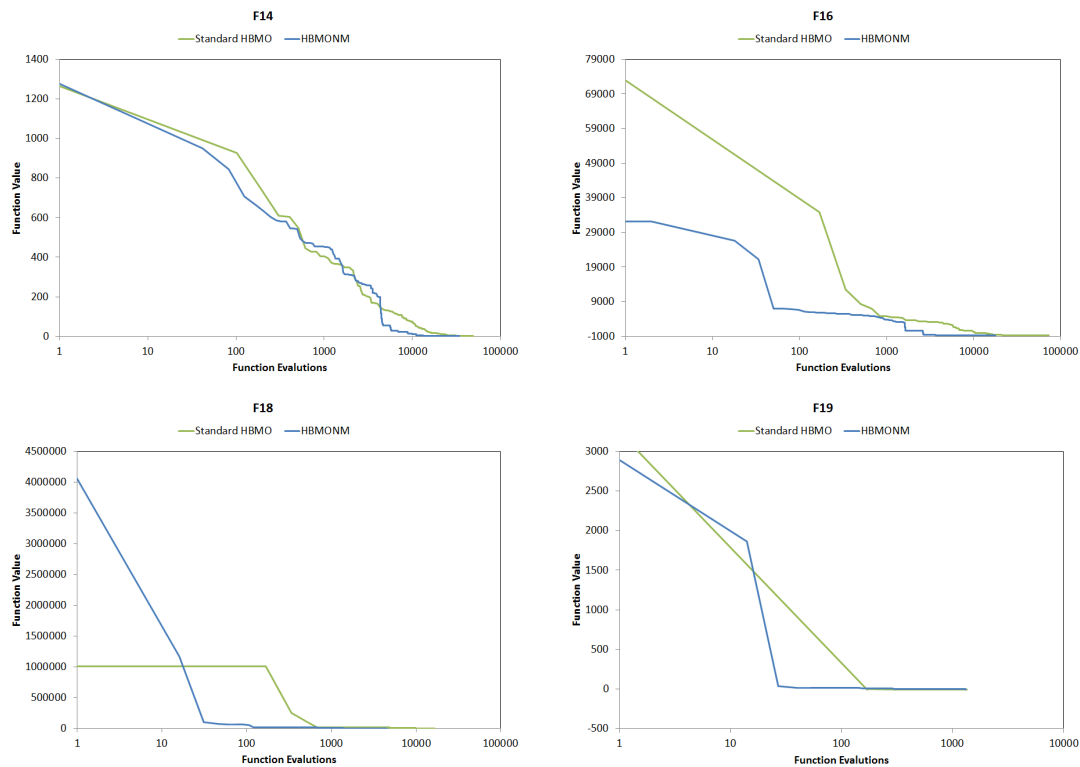


Fig. 3: The general performance of HBMONM and HBMO with integer programming problems.

integer programming problems. Further investigation may be required to improve the performance of HBMONM and HBMO on integer problems.

Acknowledgments

The research of the first author is supported in part by the Natural Sciences and Engineering Research Council of Canada (NSERC).

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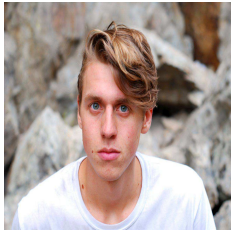
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